

*Kallson*

P#21



1653

## **RAW SEQUENCE LISTING**

PATENT APPLICATION: US/09/445,362B

DATE: 04/12/2002

TIME: 15:09:45

Input Set : A:\50125.008001.SEQLIST.TXT  
Output Set: N:\CRF3\04122002\I445362B.raw

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4 <110> APPLICANT: Hofmann, Marion Elke  
5 Domdey, Horst  
6 Henkel, Thomas  
8 <120> TITLE OF INVENTION: Myocardium-and-Skeletal  
9 Nucleic Acid, Its Preparation and Use  
12 <130> FILE REFERENCE: 50125/008001  
14 <140> CURRENT APPLICATION NUMBER: US 09/445,362B  
15 <141> CURRENT FILING DATE: 2000-05-15  
17 <150> PRIOR APPLICATION NUMBER: PCT/EP98/03584  
18 <151> PRIOR FILING DATE: 1998-06-15  
20 <150> PRIOR APPLICATION NUMBER: DE 19725186.2  
21 <151> PRIOR FILING DATE: 1997-06-13  
23 <160> NUMBER OF SEQ ID NOS: 7  
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29 <212> TYPE: DNA  
30 <213> ORGANISM: Homo sapiens  
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35 taaaatacgaa tccatcgacg aggatgaact cctcgccctcc ctgtca  
36 ggagcttagag agagagtgtgg aagacattga acctgaccgc aacctt  
37 gcaaaagagc ctgacacagaga aaacccccac agggacattc agcaat  
38 ctattggaa aaggagtccc aaaaactctt ggagaaggag aggttt  
39 ggttcagaa gacaaagagg aaagtgaaga agagcttatac ttatcc  
40 ggttctgag gaagtgtata cagaggagga ggaggaggag tcccaat  
41 agaagacagt gacgaagagg aaagaacaat taaaaactgca aaaggat  
42 aaattatgtat agtgtcaatt ctgacaactc taagccaaag atatttt  
43 gaacataaaat ttgacccaatg gcagcaatgg gaggaacaca gatgtt  
44 cccttgtgaa aatcctacag tgattgagga cgcttggac aagatcc  
45 tgacaccaca gaagtcaatt tgaacaacat tgagaacatc acaactt  
46 ctttgctgaa gccctcaagg acaacactgt ggtgaagacg ttccatc  
47 tgccgacgac agtgcagcca tggccattgc agagatgtc aaagatcc  
48 caacgtaaac gtcgagtcca acttcataac gggaaagggg atccatc  
49 ttcgcagcac aacacgggtc tcacggagct gcgtttccat aaccatc  
50 cagccaggtg gaaatggaga ttgtcaagct gctgaaggag aaccatc  
51 gggataccat tttgaactcc caggaccaag aatgagcatg acgatcc  
52 tatggataaa cagaggccaa aacgtttgca ggagcaaaaa cagcaatc  
53 aggacccaat cttaggacca aagtctggca aagaggaaca ccttagatc  
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55 gagccgtcct ctgtctcctg tggccacact teetccctct cccatc  
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59 tattctaaag gaaataaaaa attctctgag gtcagtgc当地 gagaagaaaa tggaaagacag 1620
60 ttccccac tctacccac agagatcagc tc当地gagaat ctc当地ggaag caattcgggg 1680
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62 tctttagaag aggatgcaga actgttc当地 ggtattacat gaaatgc当地 gtgagatgtt 1800
63 tctaaaatac ct当地tcaat tcaaaatgat ccctgactt当地 aaaaataatc tc当地ccatta 1860
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75 taaatac当地 gccatc当地cag aggatgaaact cctc当地ctcc ct当地cagccg aggagctgaa 180
76 ggagctagag agagaggtaa aagacattgaa acctgaccgc aacccccc当地 tggggctaaag 240
77 gcaaaagagc ct当地acagaga aaacccccc当地 agggacattc agc当地gaggagg cactgatggc 300
78 ct当地tggaa aaggagtc当地 caaaactt当地 ggagaaggag aggctgggg aatgtggaaa 360
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97 gaaaaagctc attaccagaa acattgc当地 agtcatcaaa caacaggaga gtgccaacg 1500
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104 attccaaaga gaatcttaag aaacaatcag catgtt当地 ctgtaaaat gaaaataat 1920
105 ttcttttta tgtcgtgaga tt当地gtt当地gg caagaaggc当地 ttaattt当地aa gatgtcttc当地 1980
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117 taaatacgaa tccatcgacg aggtgaact cctgcctcc ctgcagccg aggagctgaa 180  
118 ggagcttagag agagagtgg aagacattga acctgaccgc aaccttcccg tggggctaag 240  
119 gcaaaaagagc ctgacagaga aaaccccccac agggacattc agcagagagg cactgtatggc 300  
120 ctattggaa aaggagtc aaaaacttgg 121 ggagaaggag aggtgtgggg aatgtgaaaa 360  
121 ggttgcagaa gacaagagg aaagtgaaga agagcttac ttactgaaa gtaacagtga 420  
122 ggttctgag gaagtgtata cagaggagga ggaggaggag tcccaggagg aagaggagga 480  
123 agaagacagt gacgaagagg aaagaacaat tgaaactgca aaaggattt aatggactgt 540  
124 aaattatgtat agtgtcaatt ctgacaactc taagccaaag atatttaaa gtcaaataga 600  
125 gaacataaat ttgaccaatg gcaccaatgg gaggacaca gagtcggcc 126 ctgcattca 660  
126 cccttctgga aatcctacag tgattgagga cgcttggac aagattttttt gcaatgaccc 720  
127 tgacaccaca gaagtcaatt tgaacaacat tgagaacatc acaacacaga cccttacccg 780  
128 cttgctgaa gcccctcaagg acaacactgt ggtgaagacg ttcatgtctgg ccaacacgca 840  
129 tgccgacgac agtgcagcca tggccattgc agagatgtc aaagccatg agcacatcac 900  
130 caacgtaaac gtcgagtc aacttcataac gggaaagggg atcctggca tcatgagagc 960  
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132 cagccagggtg gaaatggaga ttgtcaagct gctgaaggag aacacgacgc tgctgaggct 1080  
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146 attccaaaga gaatcttaag aaacaatcag catgtttttt ctgtaaatat gaaaataaat 1920  
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156 <212> TYPE: PRT  
157 <213> ORGANISM: Homo sapiens  
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164 Leu Glu Arg Glu Leu Glu Asp Ile Glu Pro Asp Arg Asn Leu Pro Val
165           35           40          45
166 Gly Leu Arg Gln Lys Ser Leu Thr Glu Lys Thr Pro Thr Gly Thr Phe
167           50           55          60
168 Ser Arg Glu Ala Leu Met Ala Tyr Trp Glu Lys Glu Ser Gln Lys Leu
169 65           70           75          80
170 Leu Glu Lys Glu Arg Leu Gly Glu Cys Gly Lys Val Ala Glu Asp Lys
171           85           90          95
172 Glu Glu Ser Glu Glu Leu Ile Phe Thr Glu Ser Asn Ser Glu Val
173           100          105         110
174 Ser Glu Glu Val Tyr Thr Glu Glu Glu Glu Ser Gln Glu Glu
175           115          120         125
176 Glu Glu Glu Glu Asp Ser Asp Glu Glu Glu Arg Thr Ile Glu Thr Ala
177           130          135         140
178 Lys Gly Ile Asn Gly Thr Val Asn Tyr Asp Ser Val Asn Ser Asp Asn
179 145           150          155         160
180 Ser Lys Pro Lys Ile Phe Lys Ser Gln Ile Glu Asn Ile Asn Leu Thr
181           165          170         175
182 Asn Gly Ser Asn Gly Arg Asn Thr Glu Ser Pro Ala Ala Ile His Pro
183           180          185         190
184 Cys Gly Asn Pro Thr Val Ile Glu Asp Ala Leu Asp Lys Ile Lys Ser
185           195          200         205
186 Asn Asp Pro Asp Thr Thr Glu Val Asn Leu Asn Asn Ile Glu Asn Ile
187           210          215         220
188 Thr Thr Gln Thr Leu Thr Arg Phe Ala Glu Ala Leu Lys Asp Asn Thr
189 225           230          235         240
190 Val Val Lys Thr Phe Ser Leu Ala Asn Thr His Ala Asp Asp Ser Ala
191           245          250         255
192 Ala Met Ala Ile Ala Glu Met Leu Lys Ala Asn Glu His Ile Thr Asn
193           260          265         270
194 Val Asn Val Glu Ser Asn Phe Ile Thr Gly Lys Gly Ile Leu Ala Ile
195           275          280         285
196 Met Arg Ala Leu Gln His Asn Thr Val Leu Thr Glu Leu Arg Phe His
197           290          295         300
198 Asn Gln Arg His Ile Met Gly Ser Gln Val Glu Met Glu Ile Val Lys
199 305           310          315         320
200 Leu Leu Lys Glu Asn Thr Thr Leu Leu Arg Leu Gly Tyr His Phe Glu
201           325          330         335
202 Leu Pro Gly Pro Arg Met Ser Met Thr Ser Ile Leu Thr Arg Asn Met
203           340          345         350
204 Asp Lys Gln Arg Gln Lys Arg Leu Gln Glu Gln Lys Gln Glu Gly
205           355          360         365
206 Tyr Asp Gly Gly Pro Asn Leu Arg Thr Lys Val Trp Gln Arg Gly Thr
207           370          375         380
208 Pro Ser Ser Ser Pro Tyr Val Ser Pro Arg His Ser Pro Trp Ser Ser
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210 Pro Lys Leu Pro Lys Lys Val Gln Thr Val Arg Ser Arg Pro Leu Ser  
 211 405 410 415  
 212 Pro Val Ala Thr Leu Pro  
 213 420 425 430  
 214 Pro Ser Ser Gln Arg Leu Pro  
 215 435 440 445  
 216 Leu Pro Glu Lys Lys Leu Ile Thr Arg Asn Ile Ala Glu Val Ile Lys  
 217 450 455 460  
 218 Gln Gln Glu Ser Ala Gln Arg Ala Leu Gln Asn Gly Gln Lys Lys Lys  
 219 465 470 475 480  
 220 Lys Gly Lys Lys Val Lys Lys Gln Pro Asn Ser Ile Leu Lys Glu Ile  
 221 485 490 495  
 222 Lys Asn Ser Leu Arg Ser Val Gln Glu Lys Lys Met Glu Asp Ser Ser  
 223 500 505 510  
 224 Arg Pro Ser Thr Pro Gln Arg Ser Ala His Glu Asn Leu Met Glu Ala  
 225 515 520 525  
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 229 545 550  
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 235 <213> ORGANISM: Homo sapiens  
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 240 <210> SEQ ID NO: 6  
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 242 <212> TYPE: DNA  
 243 <213> ORGANISM: Homo sapiens  
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 246 <221> NAME/KEY: misc\_feature  
 247 <222> LOCATION: 12, 50, 222, 243, 255, 265  
 248 <223> OTHER INFORMATION: n = A,T,C or G  
 250 <400> SEQUENCE: 6  
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 253 atcatgagag ctctccagca caacacggtg ctcacggagc tgcggtttca taaccagagg 180  
 W--> 254 cacatcatgg gcagccaggt ggaaatggag attgtcaagc tnctgaagga gaacacgacg 240  
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 259 <212> TYPE: PRT  
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 266 20 25 30